AMENDMENTS

IN THE SPECIFICATION:

On page 10, line 2:

Primer 5T4-1: TCTTCGCCTCTTGTTGGC (gene location exon 2, 5T4 gene; Genbank accession #HSA012159; SEQ ID No. 1)

On page 10, line 4:

Primer 5T4-2: TGCAGGAAGGAACGGGA (gene location exon 1, 5T4 gene; Genbank accession #HSA012159;SEQ ID No. 2)

On page 10, line 6:

Primer 5T4-3: TTGGTAGGGAAGGAATTGGG (gene location exon 1, 5T4 gene; Genbank accession #HSA012159;SEQ ID No. 3)

On page 10, line 16:

In a preferred embodiment, 5T4 RNA is harvested approximately 1.75 milliliter aliquots of serum or plasma, with RNA extracted using the Perfect RNA Total RNA Isolation Kit (Five Prime – Three Prime, Inc., Boulder, Colorado) according to manufacturer's directions, and 10 microliters of the extracted RNA are then reverse transcribed to its cDNA as described above. Polymerase chain reaction (RT-PCR) for the 5T4 cDNA is performed using 5 microliters of the 5T4 cDNA in a final volume of 50 microliters. The reaction mixture contains one unit of Amplitaq Gold (Perkin Elmer Corp., Foster City, CA), 1 X reaction buffer, 1.5 mM $MgCl_2$, 0.2 ~~2~~ mM dNTPs, and 10 picomoles each of Primer 5T4-1 and Primer 5T4-2. The mixture is then amplified in a single-stage reaction in a thermocycler under parameters consisting of an initial 10 minute incubation at 95 degrees C, followed by about 45 cycles of denaturation at 94

On page 18, line 11:

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12. Myers *et al.*, 1994, *J. Biol. Chem.* 269: 9319-9324

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